Fri Jan 31 10:56:50 2003

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

| | 7 53 75.7 8 53 75.7 | | 2 53.5 76.4 3 53.5 76.4 | | Result Query No. Score Match Length DB |
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| 28 28 | 288 | 116 | 27 27 | 14 | gth |
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| AAB60530 AAE19032 | AAB60508 AAB60509 | AAB60516 AAB60517 | AAB60514 AAB60515 | AAU76321 | ID |
| Dog ghrelin-like G Human ghrelin pept | Rat ghrelin, SEQ I Human ghrelin, SEQ | Rat des-Gln14-ghre Human des-Gln14-gh | Rat des-Gln14-ghre Human des-Gln14-gh | Synthetic Ghrelin | Description |

| PI XX | X R | ¥ P | X B : | X PS | XX FF FF FF XX XX XX XX XX XX XX XX XX X | | ΧB | X | ž či | RES AAU ID | | | | | | | | | | | | | | | | | | | |
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|) ZE ghi 002- | -JUL-2000; | UL-2001 | -JAN-2002 | 0020825 | fi i | יו נ | C | 21-MAY-2002 | 6321; | 6321 st | | 4 4 4 4 9 9 | 49 9 | 49 | 4 4 9 9 | 49 | 4 4 ¢ | 4 4 | 44 | ٠, | 4.49 0.50 | នួន | 55 | ភូមិ | ខ្លួ | 53 53 | 53 0 | , ω ι 3 ω ι | 5 5 3 |
| ZENTARIS A i R; 2-206079/2 | 2000 | 1; 2001WO- | | 50-A2. | te 1 | y; osteopa | Ghrelin | (fir | | tandard; | | 70.0 70.0 62.9 | | | | | | | | | | | | | | | | | |
| AG. | -2201 | WO-EP07 | | | Location/Quali 3 /label= OTHER /note= "Other= | agonist; growth osteopathic. | variant | st entr | | ; peptide; | | 28 118 23 | 28 28 | 28 28 | 28 28 | 280 |) 22 K | 22 6 | 221 | 28 | 127 27 | 118 | 118 | 117 117 | 117 117 | 117 117 | 117 | 117 | 28 90 |
| | 78₽. | 929. | | | /Quali OTHER Other= | | | λ, | | | | 22 22 23 | 23 23 | 23 23 | 23 33 | 23 | 226 | 22.5 | 223 | 2 2 2 | 222 | 2 2 2 | 221 | 23 | 2 2 | 22 22 | 22 | 201 | 23 23 |
| | | | | | lifiers R r= Octanoyl Ser" | normone release; octanoyi | | | | 14 AA. | ALIGNMENTS | AAE19040 AAB60520 AAB60528 | AAE19038 AAE19039 | AAE19036 AAE19037 | AAE19034 AAE19035 | AAE19033 | AAE19029 AAE19030 | AAE19028 | AAE19021 | AAB60518 | AAB60519 | AAB65231 | AAY66708 AAU12392 | AAE23838 AAE15883 | AAB60510 AAB60511 | AAB62649 AAB20101 | AAM38890 | AAW87991 | AAE19041 ABP08975 |
| | | | | | | noy1 escer; cumour; | | | | | | | Human ghrelin pept Human ghrelin pept | Human ghrelin pept Human ghrelin pept | ghrelin ghrelin | ghrelin | ghrelin | ghrelin | an ghrelin | ine ghreli | des-GI | PRO1066 | Membrane-bound pro | Human zsig33 prote Human zsig33 prote | Rat ghrelin prepro Human ghrelin prep | ഥമ | Human polypeptide | Protein designated | Human ghrelin pept Human ORFX protein |

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RESULT 2
AAB60514
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Matches 13
      The invention relates to a novel peptide compound or its salt which induces the secretion of growth hormone and/or elevates calcium ion concentration in cells. The peptides are ghrelin homologues and are characterised in that at least one amino acid has been substituted by a modified amino acid and/or a non-amino acid compound. The invention also encompasses the unmodified peptides, the DNA encoding the peptide also encompasses the unmodified peptides.
                                                                                                                                         New peptide compounds which induce g elevate cell calcium concentrations,
                                                                                                                               of infant growth disorders
                                                                                                          Claim
                                                                                                                                                                                    WPI; 2001-159704/16.
                                                                                                                                                                                                            Kangawa K,
                                                                                                                                                                                                                                                              26-APR-2000; 2000JP-0126623
                                                                                                                                                                                                                                                                           23-JUL-1999;
29-NOV-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Growth hormone secretagogue;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a novel peptide which antagonises the effect of ghrelins when administered to a mammal. The peptide is a synthetic analogue of phrelin. Ghrelin is a 27-28 residue peptide isolated from distinctive cell types in the stomach of rate and humans and has an octancyl ester attached to a serine residue. Ghrelin is a potent releaser of growth hormone. The peptides are useful for normalising elevated growth hormone levels in mammals such as those suffering from a tumour related to overproduction of growth hormone or acromegaly. The present sequence is a ghrelin antagonising peptide of the
                                                                                                                                                                                                                                      (KANG/) KANGAWA K.
                                                                                                                                                                                                                                                                                                                                                                                           Rattus norvegicus.
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                                                                                                                                                                                                                                                                                                                                                                                                                    hormone deficiency.
                                                                                                                                                                                                                                                                                                                                                                                                                               normone secretagogue; GHS; ghrelin; concentration elevation; infant gr
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s, useful in tre
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    The invention relates to a novel peptide compound or its salt which induces the secretion of growth hormone and/or elevates calcium ion concentration in cells. The peptides are ghrelin homologues and are characterised in that at least one amino acid has been substituted by a modified amino acid and/or a non-amino acid compound. The invention also encompasses the unmodified peptides; the DNA encoding the peptides, vectors and host cells comprising such DNA; a method of producing the peptides comprising recombinant production, optionally followed by chemical modification; an antibody specific for a peptide of the invention; and an assay and kit for detecting the peptides. The peptides caused by a deficiency in growth hormone expression or activity. In particular, they are useful for promoting infant growth due to growth hormone afficiency who commoning of the invention are fully for the peptides caused by a deficiency in growth hormone expression or activity. In particular, they are useful for promoting infant growth due to growth hormone approach due to growth hormone activity.
                                                                                                                                                                                                                                                                              New peptide compounds which induce growth elevate cell calcium concentrations, usefu
                                                                                                                                                                                                                                                                                                                                                                                                                           23-JUL-1999; 99JP-0210002
29-NOV-1999; 99JP-0338841
26-APR-2000; 2000JP-0126623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    peptides comprising recombinant production, optionally followed by chemical modification; an antibody specific for a peptide of the invention; and an assay and kit for detecting the peptides. The pept of the invention are useful for treating and/or diagnosing diseases caused by a deficiency in growth hormone expression or activity. In particular, they are useful for promoting infant growth due to growth hormone deficiency. The compounds of the invention are safe with no accompanying side effects. The present sequence represents a ghrelin-type growth hormone secretagogue (GHS) of the invention.
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29-NOV-1999;
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for promoting infant growth due to growth pounds of the invention are safe with
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RESULT 4
AAB60516
CC concentration in cells. The peptides are ghrelin homologues and are characterised in that at least one amino acid has been substituted by CC a modified amino acid and/or a non-amino acid compound. The invention CC also encompasses the unmodified peptides; the DNA encoding the peptides; vectors and host cells comprising such DNA; a method of producing the comprising recombinant production, optionally followed by CC chemical modification; an antibody specific for a peptide of the CC invention; and an assay and kit for detecting the peptides. The peptides CC of the invention are useful for treating and/or diagnosing diseases CC caused by a deficiency in growth hormone expression or activity. In CC particular, they are useful for promoting infant growth due to growth CC hormone deficiency. The compounds of the invention are safe with CC no accompanying side effects. The present sequence represents a CC of the invention.
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29-NOV-1999; 99JP-0338841
26-APR-2000; 2000JP-0126623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Growth hormone secretagogue; GHS; ghrelin; precursor; preproprotein; calcium concentration elevation; infant growth disorder;
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                                                                                                                                                                                                                                                                                              Claim 3; Page 186; 210pp; Japanese
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                (KANG/) KANGAWA K.
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51.9%;
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RESULT 5
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                                                                                       The invention relates to a novel peptide compound or its salt which CC induces the secretion of growth hormone and/or elevates calcium ion concentration in cells. The peptides are ghrelin homologues and are concentration in cells. The peptides are ghrelin homologues and are concentration in cells. The peptides are ghrelin homologues and are concentration acid and/or a non-amino acid compound. The invention collaboration also encompasses the unmodified peptides; the DNA encoding the peptides; compounds the peptides; compounds of producing the peptides comprising recombinant production, optionally followed by compounds modification; an antibody specific for a peptide of the invention; and an assay and kit for detecting the peptides. The peptides of the invention are useful for treating and/or diagnosing diseases companying and effects. The promoting infant growth due to growth for mone deficiency. The compounds of the invention are safe with constant type growth hormone secretagogue (GHS) precursor protein of the invention.
Query Match
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Matches 14
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                                                      Sequence
                                                                                 ghrelin-type growth of the invention.
                                                                                                                                                                                                                                                                                                                                        Claim
                                                                                                                                                                                                                                                                                                                                                                   of infant growth disorders
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                                                      116 AA;
 Conservative
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                                                                                                                                                                                                                             The present invention provides agents for treating or preventing diseases associated with denaturation or death of neurons, which comprise a growth hormone liberation inhibitor. These can be used for treating or preventing diseases associated with denaturation or death of neurons including those due to cerebral ischaemic disorders such as cerebral infarction or oedema. Other causes of denaturation or death of neurons included Alzheimer's disease, Pick's disease, Pick's disease, Pick's disease, Pick's disease, Pick's disease, amyotrophic lateral sclerosis, diabetic neuropathy and anticancer treatments. The present sequence is a peptide described in
  AAB60508;
                        AAB60508 standard; peptide; 28
                                                                                                                                                                                                                                                                                                                                                               Disclosure;
                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                     Agents for treating diseases associated with denaturation neurons comprise growth hormone liberation inhibitor -
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Modified-site
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Pick's disease; dementia; amyotrophic lateral sclerosis; cancer;
diabetic neuropathy; neuroprotective; antlinflammatory; nootropic;
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                                                                                                                                                                                                                      exemplification of the invention.
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                                                                                                                                                                                                                                                                                                                                                             Page 17;
                                                                                                                                           Conservative
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ntia; amyotrophic lateral sclerosis; cancer;
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Best Local :
Growth hormone secretagogue; GHS; ghrelin; calcium concentration elevation; infant growth disorder;
                                        Human ghrelin,
                                                                      24-APR-2001
                                                                                                                         AAB60509 standard; peptide;
                                                                                                                                                                                                                                                                                                       Sequence
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29-NOV-1999; 99JP-0338841.
26-APR-2000; 2000JP-0126623.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   growth disorder;
                                                                                                                                                                                                                                                             .044;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Minamitake
                                                                                                                                                                                                                                                0
                                                                                                                                                                                                                                                                        Length 28
                                                                                                                                                                                                                                                Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    and
                                                                                                                                                                                                                                                14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  diagnosis
                                                                                                                                                                                                                                                Gaps
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1:

growth hormone deficiency.

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                                                                                                                                                                                                                                                 AAB60530
                                                                                                                                                                                                                                                               RESULT 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to a novel peptide compound or its salt which conditions the secretion of growth hormone and/or elevates calcium ion concentration in cells. The peptides are ghrelin homologues and are characterised in that at least one amino acid has been substituted by a modified amino acid and/or a non-amino acid compound. The invention also encompasses the unmodified peptides; the DNA encoding the peptides; vectors and host cells comprising such DNA; a method of producing the peptides comprising recombinant production, optionally followed by chemical modification; an antibody specific for a peptide of the invention; and an assay and kit for detecting the peptides. The peptides of the invention are useful for treating and/or diagnosing diseases caused by a deficiency in growth hormone expression or activity. In particular, they are useful for promoting infant growth due to growth Compone deficiency. The compounds of the invention are safe with con accompanying side effects. The present sequence represents a CC ghrelin-type growth hormone secretagogue (GHS) of the invention.
                                                                                                                                                                                                                                                                                                                                                                   Matches 14;
                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local 9
                                                                                 Growth hormone secretagogue; GHS; ghrelin; calcium concentration elevation; infant growth disorder; growth hormone deficiency.
                                                                                                                                        Dog ghrelin-like GH secretagogue peptide, SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23-JUL-1999; 99JP-0210002
29-NOV-1999; 99JP-0338841
26-APR-2000; 2000JP-0126623
                          WO200107475-A1
                                                      Canis familiaris.
                                                                                                                                                                          24-APR-2001
                                                                                                                                                                                                       AAB60530,
                                                                                                                                                                                                                                 AAB60530 standard; peptide; 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 3; Page 181; 210pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             of infant growth disorders -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New peptide compounds which induce g elevate cell calcium concentrations,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-159704/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Kangawa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (KANG/) KANGAWA K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24-JUL-2000; 2000WO-JP04907
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
                                                                                                                                                                                                                                                                                                                           1 GSSFLSPE------AKLQPR 14
                                                                                                                                                                                                                                                                                                        GSSFLSPEHORVOORKESKKPPAKLOPR 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ~
                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                               28
                                                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                                                        (first entry)
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99JP-0338841.
                                                                                                                                                                                                                                                                                                                                                                               75.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hosoda H,
                                                                                                                                                                                                                                                                                                                                                                 ; Score 53; DB;
; Pred. No. 0.04
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matsuo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         growth hormone secretion and s, useful in treatment and diagnosis
                                                                                                                                                                                                                                                                                                                                                                                                DB 22;
                                                                                                                                                                                                                                                                                                                                                                                 .044;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Minamitake
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                                                                                                                                                                                                                                                                                                                                                                                               Length 28
                                                                                                                                            NO:31.
                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                   14; Gaps
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RESULT 10
AAE19032
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             induces the secretion of growth hormone and/or elevates calcium ion concentration in cells. The peptides are ghrelin homologues and are characterised in that at least one amino acid has been substituted by a modified amino acid and/or a non-amino acid compound. The invention calso encompasses the unmodified peptides; the DNA encoding the peptides; vectors and host cells comprising such DNA; a method of producing the peptides comprising recombinant production, optionally followed by chemical modification; an antibody specific for a peptide of the invention; and an assay and kit for detecting the peptides. The peptides of the invention are useful for treating and/or diagnosing diseases considered by a deficiency in growth hormone expression or activity. In compounds of the invention are useful for promoting infant growth due to growth for mone deficiency. The compounds of the invention are safe with conscious deficiency. The compounds of the invention are safe with present sequence represents a great growth hormone secretagogue (GHS) of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Sin
Matches 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                Human; ghrelin analogue; growth-hormone secretagogue; GHS receptor; AIDS; acquired immune deficiency syndrome; weight gain; chemotherapy; dialysis; growth hormone; muscle mass; bone density; sexual dysfunction; anorexia; wasting; radiation therapy; obesity; diabetes; retinopathy; hypertension; cardiovascular disorder; gall stone; osteoarthritis; cancer; cytostatic; metabolic; immunomodulator; anti-HIV; anorectic; ophthalmological;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                           21-MAY-2002
                                                                                                                                                                                                                                                                                                                                                             AAE19032;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a novel peptide compound or its salt which induces the secretion of growth hormone and/or elevates calcium ion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  of infant growth disorders -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New peptide compounds which induce growth hormone secretion and elevate cell calcium concentrations, useful in treatment and diagnosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23-JUL-1999;
29-NOV-1999;
                                                                                                                                                   cardiant; litholytic; hepatotropic.
                                                                                                                                                                                                                                                                                       Human ghrelin peptide analogue, compound
                                                                                                                                                                                                                                                                                                                                                                                           AAE19032 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-159704/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kangawa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-APR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             24-JUL-2000; 2000WO-JP04907
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (KANG/) KANGAWA K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 μ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 GSSFLSPE-----AKLOPR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GSSFLSPEHQKLQQRKESKKPPAKLQPR 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Page 197; 210pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      28 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kojima M,
                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2000JP-0126623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           99JP-0210002.
99JP-0338841.
                                                                                                                                                                                                                                                                                                                                                                                         peptide; 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 75.7%;
50.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hosoda H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 53;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                               ₿
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    .Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 22;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .044;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  <u>,,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         K
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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25-MAY-2001; 2001WO-US17026.

06-DEC-2001. WO200192292-A2 Homo sapiens.

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RESULT 11
AAE19041
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to a truncated ghrelin analogue or their CC salt, active at growth-hormone secretagogue (GHS) receptor. Ghrelin CC analogue is useful for screening a compound capable of binding to GHS receptor and for stimulating growth hormone secretion. Ghrelin agonist is utilised for treating a growth hormone deficient state, increasing comuscle mass and bone density, treating sexual dysfunction in males or complete and search physical functioning, recovery of physical function, and/or appetite increase, or appetite increase is particularly useful for a patient complete and disease or disorder, or under going a treatment, accompanied by eight loss such as anorexia, bulimia, cancer cachexia, acquired immune deficiency syndrome (AIDS), wasting, cachexia and wasting in frail celderly and examples of treatments accompanied by eight loss include complications of diabetes weight maintenance of permanent immobilisation complications of diabetes including retinopathy, and/or cardiovascular complications of diabetes including retinopathy, and/or cardiovascular complications of diabetes including retinopathy, and/or cardiovascular diseases including hypertension, diabetes, dyslipidemias, cardiovascular complications of such diseases and for treating such diseases. Ghrelin canalogue induces growth hormone release from primary-culture pituitary cells in a dose-dependent manner without stimulating the release of other complicatory hormones. Unlike longer length dispates in analogue can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local
    Homo
                                             Human; ghrelin analogue; growth-hormone secretagogue; GHS receptor; AIDS; acquired immune deficiency syndrome; weight gain; chemotherapy; dialysis; growth hormone; muscle mass; bone density; sexual dysfunction; anorexia; wasting; radiation therapy; obesity; diabetes; retinopathy; hypertension; cardiovascular disorder; gall stone; osteoarthritis; cancer; cytostatic; metabolic; immunomodulator; anti-HIV; anorectic; ophthalmological;
                                       cardiant; litholytic;
                                                                                                                                                                                                                                          21-MAY-2002
                                                                                                                                                                                                  Human ghrelin
                                                                                                                                                                                                                                                                                                                     AAE19041 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence is human ghrelin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 4; Page 34; 37pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Truncated ghrelin analogs active at growth-hormone secretagogue receptor useful for diagnosing or treating diseases such as anorexia, bulinia, cancer, obesity, diabetes mellitus, hypertension, osteoarthritis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2002-195531/25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-MAY-2000; 2000US-207920P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (MERI ) MERCK & CO INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ituitary hormones. Unlike longer length ghrelin, ghrelin analogue can be ynthesised easily and has increased solubility in physiological buffers. he present sequence is human ghrelin peptide analogue.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                                                                                                                                       بر
                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 GSSFLSPE-----AKLOPR 14
  Bapiens
                                                                                                                                                                                                                                                                                                                                                                                                                     GSSFLSPEHORVOORKESKKPPAKLOPR 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                              peptide analogue,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ₽
K
                                                                                                                                                                                                                                                                                                                     peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     75.7%;
                                     hepatotropic
                                                                                                                                                                                                                                                                                                                       28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 53; DB
Pred. No. 0.04
0; Mismatches
                                                                                                                                                                                                                                                                                                                     B
                                                                                                                                                                                                compound 17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 23; Length 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .044;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14;
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XXXXXX

24-JUN-2002 ABP08975;

(first entry)

ABP08975 RESULT 12

ABP08975 standard;

Protein;

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The present invention relates to a truncated ghrelin analogue or their CC salt, active at growth-hormone secretagogue (GHS) receptor. Ghrelin CC analogue is useful for screening a compound capable of binding to GHS receptor and for stimulating growth hormone secretion. Ghrelin agonist is utilised for treating a growth hormone deficient state, increasing CC muscle mass and bone density treating sexual dysfunction in males or females, facilitating a weight gain, maintenance of weight, maintenance of physical functioning, recovery of physical function, and/or appetite increase, or appetite increase is particularly useful for a patient companied by a treatment, accompanied by eight loss such as anorexia, bulinia, cancer cachexia, acquired compounded efficiency syndrome (AIDS), wasting, cachexia and wasting in frail chemotherapy, radiation therapy, temporary or permanent immobilisation complications of diabetes including retinopatry, or permanent immobilisation complications of diabetes including retinopatry, and/or cardiovascular complications of diabetes including retinopathy, and/or cardiovascular complications of diabetes and certain forms of cancers, and concertain forms of cancers, and contains and certain forms of cancers, and cancers and certain forms of cancers, and certain forms of cancers and certain certain certain certain certain c
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                                                                                                                                                                                                                                                                                                XEFFXEXEXEXEXEXEXEXEFFX8X93939393939393939393939393939
                                                                                                                                                                                Best
                                                                                                                                                  Matches
                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                            Seguence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 4; Page 23; 37pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bednarek M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (MERI ) MERCK & CO INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-MAY-2001; 2001WO-US17026
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Modified-site
                                                                                                                                                                                Local
   μ
                                                                         1 GSSFLSPE----
                                                                                                                                                                                                                                                                                                                                                              present sequence
GSSFLSPEHQRVQQRKEXKKPPAKLQPR
                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                        28 AA;
                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note= "Ser(CO-(CH2)6-CH3)'
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                                                                                                                                                                                75.7%;
                                                                                                                                                                                                                                                                                                                                                          is human ghrelin peptide
                                                                     -AKLQPR 14
                                                                                                                                          ,
,
                                                                                                                                                                             Score 53;
Pred. No.
                                                                                                                                          Mismatches
                                                                                                                                                                                0
                                                                                                                                                                                DB 23
0.044;
                                                                                                                                                                                                           23; Length 28;
                                                                                                                                      o
;
                                                                                                                                                                                                                                                                                                                                                              analogue
                                                                                                                                          Indels
                                                                                                                                      14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             anorexia,
                                                                                                                                      Gaps
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                                                                                                                                               Matches
                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel human polypeptides and polynucleotides useful for diagnosing, preventing and treating cardiovascular disease, neurodegenerative, hyperproliferative disorders and autoimmune disorders -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cardiovascular disease; diabetes mellitus; systemic lupus erythematosus; hypertension; hypothyroidism; cholesterol ester storage disease; immune deficiency; immune disorder; infectious disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; open reading frame; ORFX; gene therapy; cancer; cirrhosis; hyperproliferative disorder; psoriasis; benign tumour; haemorrhage; degenerative disorder; osteoarthritis; neurodegenerative disorder;
                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        reperfusion injury in various tissues and conditions resulting from systemic cytokine damage.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-106308/14.
N-PSDB; ABN24727.
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29-AUG-2000; 2000US-228716P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    myasthenia gravis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; SEQ ID 17932; 1037pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Shimket's RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-MAY-2001; 2001WO-US10836.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200192523-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human ORFX protein sequence SEQ ID NO:17932.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (CURA-) CURAGEN CORP.
                                                                                                                                                                                     Local
   24
                                                    1 GSSFLSPE------AKLQPR 14
                                                                                                                                                                                                                                                                                                                                                           ftp.wipo.int/pub/published_pct_sequences.
GSSFLSPEHQKAQQRKESKKPPAKLQPR
                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                              90
                                                                                                                                               Conservative
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                                                                                                                                                                             75.7%;
50.0%;
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                                                                                                                                                                                 Score 53; I
   51
                                                                                                                                                                                                                   DB 23;
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RESULT 13

AAY87236 ID AAY8 XX

AAY87236 standard; Protein; 117

RESULT 14

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AAW87991
ID AAW8
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                                                                                                                                                       The present sequence represents a protein designated Zsig33. The nucleic caids are strongly expressed in stomach tissue. The polypeptide (or allelic variants/orthologs) can be used to stimulate gastric motility, can be used to stimulate gastric motility, can be used to stimulate gastric motility, can be used to treat disorders associated substance in mammals. The products are used to treat disorders associated with gastrointestinal cell contractility, secretion of digestive can substance in mammals. The products are used to treat disorders associated with gastrointestinal motility, recruitment of digestive enzymes/acids, gastrointestinal motility, recruitment of digestive can subscription regulation. Zsig33 polypeptides may also be important cultivation regulation. Zsig33 polypeptides may also be important convologically, since the family of gut-brain peptides to which the homologous protein motilin belongs has been associated with neurological and CNS functions. They may therefore be used e.g. to regulate satisty or treat obesity and other metabolic disorders where neurological feedback modulates nutritional absorption. They are useful to identify zsig33 agonists, antagonists and ligands and to produce antibodies.
                                                                                 Best
                                                                  Matches
                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human polypeptide having homology to motilin, zsig33 - useful to treat gastrointestinal motility disorders, obesity etc. and identify antagonists to treat gastrointestinal hypermotility
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24-MAR-1997;
24-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAW87991;
                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 13; Page 55-56; 69pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1999-070071/06.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Deisher TA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-MAR-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Zsig33; gastric motility; gastrointestinal inflammation; reflux disease; nutrient absorption regulation; obesity; metabolic disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Protein designated zsig33
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO9842840-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (ZYMO ) ZYMOGENETICS INC.
 24
                                                                                 Local
                                1 GSSFLSPE------AKLOPR 14
GSSFLSPEHQRVQQRKESKKPPAKLQPR
                                                                14;
                                                                                 Similarity
                                                                                                                                 117 AA;
                                                                  Conservative
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97US-0041102.
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24..117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note= "mature protein"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                              75.7%;
50.0%;
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Pred. No.
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 51
                                                                               0.2;
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Matches

Conservative

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(1 (including cancer) inflammation, cardiovascular, neurological, corproductive or developmental disorders, (e.g. arteriosclerosis, corribosis, psoriasis, acquired immune deficiency syndrome, anaemia, casthma, Crohn's disease, microbial or other infections, congestive or ischaemic heart disease, Alzheimer's, Parkinson's or Huntington's congestive or concleic acids can be used for the recombinant production of HSPP, for detecting HSPP in standard hybridisation and amplification assays (for criplex-forming or ribozyme therapeutics, for detecting related sequences or genetic variations, and for chromosomal mapping. HSPP are also used to raise specific antibodies (Ab) and to screen for agonists and accompletity antibodies (Ab) and to screen for agonists or antiscore, hSPP-related diseases (in usual immunoassays), as therapeutic antisponists, in competitive drug screens, and for purification of HSPP cantagonists, in competitive drug screens, and for purification of HSPP antiscore.
  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAZ98109 to AAZ98242 encode AAY87224 to AAY87357 which represent the human signal peptide-containing proteins HSPP-1 to HSPP-134. HSPPs have anticancer, anti-inflammatory, antimicrobial, nootropic, hepatotropic, neuroprotective, cardiovascular and antiasthmatic activities, and can be used in gene therapy. HSPPs can be used to treat or prevent disorder associated with decreased activity or function of HSPP. Antagonists of HSPP are used to treat or prevent disorders associated with increased activity or function of HSPP. Such diseases include cell proliferation
                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Lal P, T
Akerblom
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31-JUL-1998;
01-OCT-1998;
11-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New human signal peptide-containing proteins useful in treatment, prevention and diagnosis of e.g. cancer, inflammation and cardiovascular disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; signal peptide-containing protein; HSPP; diagnosis; cancer; inflammation; cardiovascular disease; anticancer; anti-inflammatory; antimicrobial; nootropic; neuroprotective; cardiovascular; hepatotropic; antiasthmatic; gene therapy; cell proliferation; neurological disorder; reproductive disorder; developmental disorder; arteriosclerosis; cirrhosis; psoriasis; acquired immune deficiency syndrome; anaemia; asthma; Crohn's disease; infection; Alzheimer's disease; schizophrenia; parkinson's disease; Huntington's disease; ovulatory defect;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             06-JAN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200000610-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Page 168-169;
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Au-Young
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98US-0094983.
98US-0102686.
98US-0112129.
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H, Patterson C, Reddy
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53;
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R, Hillman
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RESULT 15
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                                                                                              The invention relates to human nucleic acids (AAI57798-AAI61369) and the encoded polypeptides (AAM38642-AAM42213) with nootropic, immunosuppressant and cytostatic activity. The polynucleotides are useful in gene therapy. A composition containing a polypeptide or polynucleotide of the invention may be used to treat diseases of the peripheral nervous system, such as peripheral nervous injuries, peripheral neuropathy and localised neuropathies and central nervous system diseases, such as alzheimer's, Parkinson's disease, Huntington's disease, amyotrophic lateral sclerosis, and Shy-Drager Syndrome. Other uses include the utilisation of the activities such as: Immune system suppression, activity, chemotactic/chemokinetic activity, haemostatic and thrombolytic activity, cancer diagnosis and therapy, drug screening, assays for receptor activity, arthritis and inflammation, leukaemias and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-JUL-2000;
03-AUG-2000;
14-SEP-2000;
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25-APR-2000;
09-JUL-2000;
                                                                                    assays for receptor activity, Ca.N.S disorders.
                                                specification.
                                                                                                                                                                                                                                                                                                                                                                                                                     Example
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29-NOV-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            peripheral nervous system; neuropathy; central nervous system; CNS; Alzheimer's; Parkinson's disease; Huntington's disease; haemostatic; amyotrophic lateral sclerosis; Shy-Drager Syndrome; chemotactic; chemokinetic; thrombolytic; drug screening; arthritis; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (HYSE-)
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                                                                disorders.
The sequen
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DB; AAI58046.
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Wang :
Zhou
                                                                  sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO
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2000US-0552317.

2000US-05598042.

2000US-0620312.

2000US-0653450.

2000US-0663191.

2000US-0693036.

2000US-0693036.
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Best Local Similarity 50.0%; pred: No. 0.2;
Matches 14; Conservative 0; Mismatches 0; Indels 14; Gaps 1;
Matches 14; Conservative 14; Gaps 1;
Matches 14; Conservative 0; Mismatches 0; Indels 14; Gaps 1;
Matches 14; Conservative 0; Indels 14; Gaps 14; G
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Maximum Match 100%
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Perfect score:
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     protein search, using sw model
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2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*

3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*

4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*

5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*

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US-09-460-145-4
US-09-500-358-3
US-09-500-358-3
US-09-605-430-3
US-08-659-873-3
US-08-659-473-3
US-08-618-957A-12
US-08-618-364A-2
US-08-618-364A-2
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US-08-640-389A-12
US-08-343-365A-9
US-08-343-760A-2
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US-08-822-897C-2
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ALIGNMENTS

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Sequence 2, Application US/09046479
PALENT NO. 6291653
GENERAL INFORMATION:
APPLICANT: Sheppard, Paul O.
APPLICANT: Sheppard, Paul O.
TITLE OF INVENTION: MOTILIN HOMOLOGS
NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
ADDRESSEE: ZymoGenetics, Inc.
STREET: 1201 Eastlake Avenue East
CITY: Seattle
STATE: WA
                                                                      ; MOLECULE TYPE:
; FRAGMENT TYPE:
US-09-046-479-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 1
US-09-046-479-2
Query Match
Best Local S
Matches 14
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SEQUENCE CHARACTERISTICS:
LENGTH: 117 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/046,479
FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                     FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Sawislak, Deborah A
REGISTRATION NUMBER: 37,438
REFERENCE/DOCKET NUMBER: 97-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION:
TELEPHONE: 206-442-6672
                                                                                                                                                                                                                                  TELEX:
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                  75.7%;
Score 53; DB Pred. No. 0.01
                  DB 4;
0.015;
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Gaps

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GSSFLSPE-----AKLOPR

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APPLICANT: Sheppard, Paul O.
APPLICANT: Jaspers, Stephen R.
APPLICANT: Jaspers, Stephen R.
APPLICANT: Bishop, Paul D.
TITLE OF INVENTION: SGIP PEPTIDES
FILE REFERENCE: 99-51
CURRENT APPLICATION UMBER: US/09/608,810A
CURRENT FILING DATE: 2000-06-30
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                                                                                                                           Sequence 4, Application US/09608810A Patent No. 6420521
GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: FRAGMENT TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: 206-442-6672
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/822,897C FILING DATE: CLASSIFICATION: 536 PRIOR APPLICATION DATA: APPLICATION NUMBER: FILING DATE:
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APPLICANT: Deisher, Theresa A.
TITLE OF INVENTION: MOTILIN HOMOLOGS
NUMBER OF SEQUENCES: 7
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                                                                                                                                                                                                                                    24 GSSFLSPEHQRVQQRKESKKPPAKLQPR 51
                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 117 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Sawislak, Deborah A
REGISTRATION NUMBER: 37,438
REFERENCE/DOCKET NUMBER: 97-
                                                                                                                                                                                                                                                                   1 GSSFLSPE------AKLOPR 14
                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY:
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1201 Eastlake Avenue
                                                                                                                                                                                                                                                                                                      Conservative
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internal
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APPLICANT: Jaspers, Stephen R.
APPLICANT: Deisher, Theresa A.
APPLICANT: Deisher, Theresa A.
APPLICANT: Deisher, Theresa A.
ITILE OF INVENTION: SGIP PEPTIDES
ITILE OF INVENTION: SGIP PEPTIDES
CURRENT APPLICATION NUMBER: US/09/608,810A
CURRENT FILING DATE: 2000-06-30
PRIOR APPLICATION NUMBER: 60/141,592
PRIOR FILING DATE: 1999-06-30
NUMBER OF SEQ ID NOS: 7
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 2
LENGTH: 11
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                            RESULT 5
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Patent No. 5206161
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 60/141,592
PRIOR FILING DATE: 1999-06-30
NUMBER OF SEQ ID NOS: 7
SOFTWARE: FASTSEQ for Windows Version
SEQ ID NO 4
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Best Local (
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ZIP: 94080
COMPUTER READABLE FORM:
MEDIUM TYDE: 5.25 inch, 360 Kb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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TYPE: PRT
                                                                                                                                                           APPLICANT: Dennis Drayna and Daniel Eaton
TITLE OF INVENTION: No. 5206161el Plasma Carboxypeptidase
NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
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NAME/KEY: SIGNAL
LOCATION: (1)...
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                                                                                                  STREET: 460 Point San Bruno Blvd
CITY: South San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity nes 8; Conserv
                                                                      COUNTRY: U
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Similarity 50.0%;
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100.0%;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                     DB 4;
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INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 417 amino acids
TYPE: AMINO ACID
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:

MEDIUM TYPE: 5.25 inch, 360 Kb floppy
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Parin (Genentech)
CURRENT APPLICATION DATA:
CURRENT APPLICATION NUMBER: US/08/277,540
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                                                         CLASSIPICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 08/167727
FILING DATE: 15-DEC-1993
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 07/959944
FILING DATE: 14-OCT-1992
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 07/649591
FILING DATE: 01-FEB-91
FILING DATE: 01-FEB-91
ATTORNEY/AGENT INFORMATION:
NAME: Hasak, Janet E.
REGISTRATION UNMBER: 28,616
REGISTRATION UNMBER: 689D1C1D1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION: Dennis T., Eaton, Dan L.
APPLICANT: Drayna, Dennis T., Eaton, Dan L.
TITLE OF INVENTION: No. 5474901el Plasma Carboxypeptidase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: HABAK, JANET E.
REGISTRATION NUMBER: 28,616
REFERENCIOCKET NUMBER: 688
TELECOMMUNICATION INFORMATION:
TELECHONE: 415/266-1896
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-1896
TELEPAX: 415/952-9881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
STREET: 460 Point San Bruno Blvd
CITY: South San Francisco
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PRIOR APPLICATION DATA:
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CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                385 SGFLLPESRIKP 396
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hes 6; Conservarium
                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/01 FILING DATE: 19-JUL-1994
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Query Match
Best Local Similarity
Matches 6; Conserve
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US-08-430-787A-7
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GENERAL INFORMATION:
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FILING DATE: 27-APR-1995
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/277,540
FILING DATE: 19-UUL-1994
APPLICATION NUMBER: 08/167727
FILING DATE: 15-DEC-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/959944
FILING DATE: 14-OCT-1992
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 07/649591
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 07/649591
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 07/649591
ATTORNEY/AGENT INFORMATION:
NAME: USAGN INFORMATION:
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LENGTH: 417 amino acids
TYPE: amino acid
                                                                                                                                                                                               TELEX: 910/371-7168
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:

COMPUTER TYPE: 5.25 inch, 360 Kb f

COMPUTER: IBM PC compatible

COMPUTER: IBM PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: patin (Genen CURRENT APPLICATION DATA:
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech,
                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION: TELEPHONE: 415/225-1896
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Drayna, TITLE OF INVENTION:
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SOFTWARE: patin (
                                                                                                                                                                                                                                                                                                                                        NAME: Hasak, Janet E. REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: 460 Point San Bruno Blvd
CITY: South San Francisco
STATE: California
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                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER:
                                                                                                                                                                            LENGTH:
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2 SSFLSPEAKLOP 13
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                                         Conservative
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No. 5593674el Plasma Carboxypeptidase
                                                          52.9%;
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                                       Score 37; DB 1;
Pred. No. 57;
4; Mismatches
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                                                                          Length 417;
                                         Indels
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; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-09-460-145-2
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Best Local Similarity 7/...
Thes 7; Conservative
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                                                                                                                                                                                                                                           Sequence 4, Application US/09460145 Patent No. 6287838
                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Kriz, Ron
APPLICANT: Song, Chuanzheng
TITLE OF INVENTION: CYTOSOLIC PHOSPHOLIPASE A2-BETA ENZYMES
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 2, Application US/09460145 Patent No. 6287838 GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: Brown, Scott A.
REGISTRATION NUMBER: 32,724
REFERENCE/DOCKET NUMBER: GI5289
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 498-8224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genetics Institute, Inc.
STREET: 87 CambridgePark Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Kriz, Ron
APPLICANT: Song, Chuanzheng
TITLE OF INVENTION: CYTOSOLIC PHOSPHOLIPASE A2-BETA ENZYMES
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APPLICATION NUMBER:
                                                                             STREET: 87 Cambridge
                                                                                                                                                                                                                                                                                                                                         668 SPEEQLOPR 676
                                                      COUNTRY:
                                                                                                                         ADDRESSEE:
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USA
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                                                                                                     87 CambridgePark Drive
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Pred. No. 1.2e+02;
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             COMPUTER: IM PC COMPATIBLE
COMPUTER: IM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PALENITIA Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/827,208
FILING DATE: 28-MAR-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/014,608
FILING DATE: 29-MAR-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/041,264
FILING DATE: 19-MAR-1997
PRIOR APPLICATION NUMBER: US 60/041,264
FILING DATE: 19-MAR-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 3, Application Patent No. 6025178
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 797 amino acid
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Kramer, Ruth M.
APPLICANT: Pickard, Richard T.
APPLICANT: Sharp, John D.
APPLICANT: Strifler, Beth A.
APPLICANT: Strifler, Beth A.
TITLE OF INVENTION: HUMAN PHOSPHOLIPASE A2 AND RELATED
TITLE OF INVENTION: NUCLEIC ACID COMPOUNDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE:
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FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 498-8224
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                   STREET: Lilly Corpo
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TOPOLOGY: li
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REGISTRATION NUMBER: 32
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     6 SPEAKLOPR 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC
OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                     46285
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Lilly Corporate Center
                                                                                                                                                                                                                                                                                                                   United States of America
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chiou, Xue-Chiou C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (617) 876-5851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                              Floppy disk
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77.8%;
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Pred. No.
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Query Match
Best Local Similarity
Thes 7; Conserve
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                                                                                                                                                                       CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/827,208
FILING DATE: 28-MAR-1997
APPLICATION NUMBER: US 60/014,608
FILING DATE: 29-MAR-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/041,264
FILING DATE: 19-MAR-1997
ATTORNEY,AGENT INFORMATION:
NAME: GAY10, Paul J.
REGISTRATION NUMBER: 36,808
TEGISTRATION NUMBER: X-10610
                                                                                              REFERENCE/DOCKET NUMBER: X-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (317) 276-0756
TELEPAX: (317) 276-3861
INFORMATION FOR SEQ ID NO: 3:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (317) 276-3861 INFORMATION FOR SEQ ID NO: 3: SEQUENCE CHARACTERISTICS: LENGTH: 913 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                equence 3, Application US/09500358 atent No. 6197569
                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/500,358
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Strifler, Beth A.
TITLE OF INVENTION: HUMAN PHOSPHOLIPASE A2 AND RELATED
TITLE OF INVENTION: NUCLEIC ACID COMPOUNDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Eli Lilly and Company
STREET: Lilly Corporate Center
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                                                                             SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Kramer, Ruth M.
APPLICANT: Pickard, Richard T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       803 SPEEQLOPR 811
                            LENGTH: 913 amino acid
                        TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Indianapolis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
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TYPE: protein
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                                                       913 amino acide
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Strifler, Beth
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Pred. No. 1.4e+02;
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Query Match
Best Local Similarity
Watches 7; Conserv
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Patent No. 6242206
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/827,208
FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/041,264
                                                                                                                                                                                                       TELEPHONE: (317) 276-0756
TELEFAX: (317) 276-3861
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
SEQUENCE: CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/498,809
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Strifler, Beth A.
TITLE OF INVENTION: HUMAN PHOSPHOLIPASE A2 AND RELATED
TITLE OF INVENTION: NUCLEIC ACID COMPOUNDS
                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION: TELEPHONE: (317) 276-0756
                                                                                                                                                                                                                                                                                                                                            FILING DATE: 19-MAR-1997
ATTORNEY/AGENT INFORMATION:
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803 SPEEQLQPR 811
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               803 SPEEQLOPR 811
                                                                                                                                                           LENGTH: 913 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: Lilly Corpo
                                                                                                                                                                                                                                                                                              NAME: Gaylo, Paul J.
REGISTRATION NUMBER: 36.
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE:
                              6 SPEAKLQPR 14
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Lilly Corporate Center
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Pickard, Richard T.
Sharp, John D.
                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chiou, Xue-Chiou C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                52.9%;
77.8%;
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                                                                                                                                                                                                                                                                                                               36,808
                                                                                Score 37; DB 4;
Pred. No. 1.4e+02;
                                                                                                                                                                                                                                                                                               X-10610
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 37; DB 4;
Pred. No. 1.4e+02;
1; Mismatches 1
                                                                Mismatches
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                                                                Indels
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RESULT 13 US-08-598-873-36

Sequence 36, Application US/08598873 Patent No. 5928884 GENERAL INFORMATION:

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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/598,873
FILING DATE: 09-FEB-196
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Friebel, Thomas E.
REGISTRATION NUMBER: 29,258
REFERENCE/DOCKET NUMBER: 8666-004
TELEPHONE: (212) 790-9090
TELEPHONE: (212) 869-9741/8864
                                 COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/605,430
FILING DATE: 22-FEB-1996
CLASSIFICATION: 536
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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LENGTH: 57 amino acids
TYPE: amino acid
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Croce, Carlo M.
APPLICANT: Huebner, Kay
TITLE OF INVENTION: FHIT PROTEINS AND NUCLEIC ACIDS AND
TITLE OF INVENTION: METHODS BASED THEREON
NUMBER OF SEQUENCES: 86
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Croce, Carlo M. APPLICANT: Huebner, Kay
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CITY: New York
STATE: New Yor
                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: New York
COUNTRY: U.S.A.
ZIP: 10036-2711
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: New York
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                                                                                             ; MOLECULE TYPE: protein PCT-US91-08177-13
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GENERAL INFORMATION:
APPLICANT: Samal, Siba K
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TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                          TELEFAX: 202-962-8300 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                     APPLICATION NUMBER: PCT/US:
FILING DATE: 19911104
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/
FILING DATE: 05-NOV-1990
ATTORNEY/AGENT INFORMATION:
NAME: Highet, David W
REGISTRATION NUMBER: 30,26:
REFERENCE/DOCKET NUMBER: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REGISTRATION NUMBER: 29,258
REFERENCE/DOCKET NUMBER: 860
TELECOMMUNICATION INFORMATION:
                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 263 amino acids
TYPE: AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 19
CORRESPONDENCE ADDRESS:
ADDRESSEE: Venable, Baetjer, Howard & Civiletti
STREET: 1201 New York Avenue N.W., suite 1000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION: TELEPHONE: 202-962-4854
                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: peptide
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Local Similarity 58.3%;
nes 7; Conservative
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OPERATING SYSTEM:
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6 SPEAKLOPR 14
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SYSTEM: PC-DOS/MS-DOS
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1; Mismatches
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                                                 Score 35; DB
Pred. No. 78;
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Search completed: January 29, 2003, 13:24:20 Job time : 15 Becs

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re greater than or equal to the score of the result being printed,
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ALIGNMENTS

C;Accession: A59316
R;KOjima, M.; Hosoda, H.; Date, Y.; Nakazato, M.; Matsuo, H.; Kangawa, K.
Nature 402, 656-660, 1999
A;Title: Ghrelin is a growth-hormone-releasing acylated peptide from stomach.
A;Reference number: A59316; MUID:20067959; PMID:10604470

A; Molecule type: mRNA A; Residues: 1-117 < KOJ>

A; Accession: A59316

A;Status: not compared with conceptual translation

ghrelin precursor - human
N;Alternate names: preproghrelin
C;Species: Homo sapiens (man)
C;Date: 16-Jun-2000 #sequence_revision 16-Jun-2000 #text_change 21-Jul-2000

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ghrelin precursor - rat
N;Alternate names: preproghrelin
C;Species: Rattus norvegicus (Norway rat)
C;Date: 16-Jun-2000 #sequence_revision 16-Jun-2000 #text_change 21-Jul-2000
C;Date: 16-Jun-2000 #sequence_revision 16-Jun-2000 #text_change 21-Jul-2000
C;Accession: B59316
R;Kojima, M.; Hosoda, H.; Date, Y.; Nakazato, M.; Matsuo, H.; Kangawa, K.
Nature 402, 656-660, 1999
A;Title: Ghrelin is a growth-hormone-releasing acylated peptide from stomach
A;Title: Ghrelin is a growth-hormone-releasing acylated peptide from stomach
A;Reference number: A59316; MUID:20067959; PMID:10604470
A;Accession: B59316
A;Accession: B59316
A;Status: not compared with conceptual translation
A;Residues: 1-17 < KOJJ
A;Cross-references: GB:AB029433; NID:96691569; PIDN:BAA89370.1; PID:96691570
A;Cross-references: GB:AB029433; NID:96691569; PIDN:BAA89370.1; PID:96691570
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C; Comment: Ghrelin secreted by the stomach stimulates the C; Superfamily: motilin
C; Keywords: hormone; lipoprotein; stomach
F; 1-23/Domain: signal sequence #status predicted <SIG>F; 24-51/Product: ghrelin #status predicted <MAT>F; 52-11/Pomain: carboxyl-terminal propeptide #status predicted control properties #status p
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A; Experimental source: strain SD; tissue stomach endocrine
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Pred. No. 0.027;
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A-D-glcr

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glycoprotein G - bovine respiratory syncytial virus (isolate FS-1 C;Species: bovine respiratory syncytial virus C;Species: bovine respiratory syncytial virus C;Date: 14-Jul-1994 #sequence_revision 14-Jul-1994 #text_change C;Accession: PQ0768

R;Mallipeddi, S.K.; Samal, S.K.

J. Gen. Virol. 74, 2001-2004, 1993

A;Title: Sequence variability of the glycoprotein gene of bovine A;Reference number: JQ2284; MUID:93389461; PMID:8376974

A;Accession: PQ0768

A;Molecule type: mRNA
A;Residues: 1-250 <MAL>
C;Superfamily: respiratory syncytial virus major surface glycoprotein c;Superfamily: respiratory syncytial virus major surface glycoprotein; transmembrane protein
F;1-31/Domain: intracellular #status predicted <INT>
F;32-59/Domain: extracellular #status predicted <EXT>
F;60-250/Domain: extracellular #status predicted <EXT>
F;78,156,226,241,244/Binding site: carbohydrate (Asn) (covalent)
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C.Paccession: PQ0769
R.Mallipeddi, S.K.; Samal, S.K.
J. Gen. Virol. 74, 2001-2004, 1993
A;Title: Sequence variability of the glycoprotein gene of bov
A;Reference number: JQ2284; MUID:93389461; PMID:8376974
A;Accession: PQ0769
A;Molecule type: mRNA
A;Residues: 1-248 <MAL>
A;Experimental source: isolate VC464
A;Note: the authors translated the codon ACC for residue 85 a
C;Superfamily: respiratory syncytial virus major surface glyc
C;Keywords: glycoprotein; transmembrane protein
C;Keywords: glycoprotein; transmembrane protein
F;1-29/Domain: intracellular #status predicted <INT>
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C;Comment: Ghrelin secreted by the stomach stimulates t
C;Superfamily: motilin
C;Keywords: hormone; lipoprotein; stomach
F;1-23/Domain: signal sequence #status predicted <SIG>F;24-51/Product: ghrelin #status predicted <MAT>
F;52-117/Domain: carboxyl-terminal propeptide #status |
F;26/Binding site: octanoate (Ser) (covalent) #status |
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50.0%;
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Pred. No. 4.1;
1; Mismatches
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Pred. No. 0.02
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protein K02E2.7 -
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A; NEGLUB. F----
A; Molecule type: DNA
A;Residues: 1-105 <TIGR>
A;Cross-references: GB:U39688; GB:L43967; NID:g1045753; PID:g1045769; TIGR:MG092
                                                                                                                                                                                                                                                                                                                                                                                                 R;Fraser, C.M.; Gocayne, J.D.; White, O.; Adams, M.D.; Claytor, M.; Fuhrmann, J.; Nguyen, D.; Utterback, T.R.; Saudek, D.M.; I., C.A.; Venter, J.C.
Science 270, 397-403, 1995
A;Title: The minimal gene complement of Mycoplasma genitalium.
A;Reference number: A64200; MUID:96026346; PMID:7569993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C;Species: Mycoplasma genitalium
C;Date: 17-Nov-1995 #sequence_revision 17-Nov-1995 #text_change 07-Dec-1999
C;Accession: B64210
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C; Superfamily: Escherichia coli ribosomal
C; Keywords: protein biosynthesis; ribosome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R;Himmelreich, R.; Hilbert, H.; Plagens, H.; Pirkl, E.; Li, B.C.; Herrmann, R. Nucleic Acids Res. 24, 4420-4449, 1996
A;Title: Complete sequence analysis of the genome of the bacterium Mycoplasma A;Reference number: S73327; MUID:97105885; PMID:8948633
A;Accession: S73927
                                                                                                                                                                                                                                                                                                                                                     A;Status: preliminary; nucleic acid sequence not shown; translation
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A; Residues: 1-104 <HIM>
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N.Alternate names: hypothetical protein G07_orf104b
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                                                                                                           Score 42;
Pred. No. 2
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Pred. No. 2
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                                                                                                                              Length 105;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Clayton, R.A.; Fleischmann, R. D.M.; Phillips, C.A.; Merrick,
                                                                                      Indels
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Caenorhabditis elegans

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conserved hypothetical protein CC0691 [imported] - Caulobacter crescentus C;Species: Caulobacter crescentus C;Species: Caulobacter crescentus C;Date: 20-Apr-2001 #sequence_revision 20-Apr-2001 #text_change 20-Apr-2001 C;Accession: H8734
R;Nierman, W.C.; Feldblyum, T.V.; Paulsen, I.T.; Nelson, K.B.; Bisen, J.; Heidelberg, J. B.; Laub, M.T.; DeBoy, R.T.; Dodson, R.J.; Durkin, A.S.; Gwinn, M.L.; Haft, D.H.; Kolon, J.; Ermolaeva, M.; White, O.; Salzberg, S.L.; Shapiro, L.; Venter, J.C.; Fraser, C.M. Proc. Natl. Acad. Sci. U.S.A. 98, 4136-4141, 2001
A;Title: Complete Genome Sequence of Caulobacter crescentus.
A;Reference number: A87249; MUID:21173698; PMID:11259647
A;Accession: H87334
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Accession: S34851
A;Status: preliminary; translation not A;Molecule type: DNA
A;Residues: 1-283 <FEL>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               R;Felsberg, J.; Petricek, M.; Tichy, P.
Nucleic Acids Res. 21, 3582, 1993
A;Title: Nucleotide sequence of the mini-plasmid pSLG33
A;Reference number: S34850; MUID:93348001; PMID:8346038
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     hypothetical 31.9K protein - Streptomyces lavendulae plasmid pSLG33 C;Species: Streptomyces lavendulae C;Date: 19-Mar-1997 #sequence_revision 25-Apr-1997 #text_change 20-1C;Accession: S34851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT
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A;Reference number: Z19716
A;Accession: T23249
A;Cross-references: GB:AE005673; NID:gl3421912; PIDN:AAK22676.1; GSPDB:GN00148 C;Genetics:
A;Gene: CC0691
                                                               A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-208 <STO>
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C;Superfamily: Streptomyces lavendulae plasmid pSLG33 hypothetical 31.9K protein
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Best Local (
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Best Local (
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Experimental source: clone K02E2
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Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 15-Oct-1999
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nes 8; Conserv
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Query Match
Best Local Similarity
Marches 8; Conserve
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A; Authors: Kreft, J.; Kuhn, M.; Kunst, F.; Kurapkat, G.; Madueno, E.; ok, C.; Schlueter, T.; Simoes, N.; Tierrez, A.; Vazquez-Boland, J.A.; A;Title: Comparative genomics of Listeria species.

A;Reference number: AB1077; MUID:21537279; PMID:11679669
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 R;Nierman, W.C.; Feldblyum, T.V.; Paulsen, I.T.; Nelson, K.E.; Eisen, J.; Heidelberg, B.; Laub, M.T.; DeBoy, R.T.; Dodson, R.J.; Durkin, A.S.; Gwinn, M.L.; Haft, D.H.; Koln, J.; Ermolaeva, M.; White, O.; Salzberg, S.L.; Shapiro, L.; Venter, J.C.; Fraser, C. Proc. Natl. Acad. Sci. U.S.A. 98, 4136-4141, 2001

A;Title: Complete Genome Sequence of Caulobacter crescentus.

A;Reference number: A87249; MUID:21173698; PMID:11259647
                                                                                                                                                                                                                                                                                                                                                                                                                                C;Accession: AB1086
R;Glaser, P.; Frangeul, L.; Buchrieser, C.; Amend, .; Dominguez-Bernal, G.; Duchaud, E.; Durand, L.;
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                                                                                                                                                                                                                                 A; Molecule type: DNA
A; Residues: 1-343 <GLA>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C;Species: Listeria monocytogenes
C;Date: 27-Nov-2001 #sequence_revision 27-Nov-2001 #text_change 27-Nov-2001
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C;Species: Caulobacter crescentus
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A; Residues: 1-308 <STO>
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                                    2 SSFLSPEAKLQ 12
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7; Conserv
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Pred. No. 18;
2; Mismatches
                                                                                              Score 40;
Pred. No.
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. 18;
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L.; Dussurget, O.;
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Fsihi, H
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RESULT 12 AH1449

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A; Molecule type: DNA
A; Acidecule type: DNA
A; Residues: 1-709 MUR>
A; Cross-references: EMBL: Z50046; NID: g899393; PIDN: CAA90369.1; PID: g899396; GSPDB: GN0000
A; Experimental source: strain AB972
A; Experimental source: A66200; PDB: INCS
A; Experimental source: A66200; PDB: INCS
A; Reference number: A66200; PDB: INCS
A; Reference number: A67819; PDB: ISFD
A; Contents: annotation; conformation by (1)H-NMR, residues 577-608
A; Reference number: A67819; PDB: ISFD
A; Contents: annotation; conformation by (1)H-NMR, residues 577-608
A; Nol. Biol. 228, 619-636, 1992
A; Title: Adjacent zinc-finger motifs in multiple zinc-finger peptides from SWI5 form str
A; Reference number: A58624; MUID: 93085740; PMID: 1453467
A; Reference number: A58624; MUID: 93085740; PMID: 1453467
A; Reference number: A58624; MUID: 93085740; PMID: 1453467
A; Rouhaus, D.; Nakaseko, Y.; Schwabe, J.W.R.; Klug, A.
J. Mol. Biol. 228, 637-651, 1992
A. Title: Af8624; MUID: 93085740; PMID: 1453467
A; Rouhaus, D.; Nakaseko, Y.; Schwabe, J.W.R.; Klug, A.
J. Mol. Biol. 228, 637-651, 1992
A. Title: Af8624; MUID: 93085740; PMID: 1453467
A; Rouhaus, D.; Nakaseko, Y.; Schwabe, J.W.R.; Klug, A.
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A;Residues: 1-709 <STI>A;Cross-references: EMBL:X06978; NID:g4595; PIDN:CAA30040.1;
A;Cross-references: EMBL:X06978; NID:g4595; PIDN:CAA30040.1;
A;Murphy, L.; Richards, C.; Harris, D.
submitted to the EMBL Data Library, July 1995
A;Reference number: S57971
A;Accession: S57973
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  transcription factor SWI5 [validated] - yeast (Saccharomyces cerevisiae) N;Alternate names: protein YD8358.03c; protein YDR146c C;Species: Saccharomyces cerevisiae C;Date: 30-Jun-1992 #sequence revision 30-Jun-1992 #text_change 08-Dec-2000 C;Accession: S00342; S57973
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Science 294, 849-852, 2001
A;Authors: Kreft, J.; Kuhn, M.; Kunst, F.; Kurapkat, G.; Madueno, E.; ok, C.; Schlueter, T.; Simoes, N.; Tierrez, A.; Vazquez-Boland, J.A.; A;Title: Comparative genomics of Listeria species.
A;Reference number: AB1077; MUID:21337279; PMID:11679669
A;Accession: AH1449
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A;Experimental source: strain Clip11262
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                                                    Reference number: A58623; MUID:93085741; PMID:1453468
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MBO J. 7, 485-494, 1988
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8; Conserv
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72.7%;
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20;
          which codes
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Voss, H.; W
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Fsihi, H.
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A;Accession: T49100
                                                                                                                                                                                                                                                                                                                             hypothetical protein F4F15.320 - Arabidopsis thaliana C;Species: Arabidopsis thaliana (mouse-ear cress) C;Species: 02-Jun-2000 #sequence_revision 02-Jun-2000 #text_change 02-Jun-2000
     A; Map posit
A; Introns:
                                                                      C; Genetics:
                                                                                                  A; Experimental source: cultivar
                                                                                                                                            A; Molecule type: DNA
A; Residues: 1-371 < ALC>
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A;Reference number: A36408; MUID:91012801;
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C;Superfamily: transcription factor SWI5
C;Keywords: DNA binding; metalloprotein; transcription fac
F;550-574/Region: zinc finger CCHH motif
F;580-604/Region: zinc finger CCHH motif
F;580-632/Region: zinc finger CCHC motif
F;609-632/Region: zinc finger CCHC motif
F;552,557,570,574/Binding site: zinc (Cys, Cys, His, His)
F;582,887,600,604/Binding site: zinc (Cys, Cys, His, His)
F;611,613,626,632/Binding site: zinc (Cys, Cys, His, Cys)
                                                  A; Gene: ATSP: F4F15.320
                                                                                                                      A; Cross-references:
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J. Virol. 64, 5559-5569, 1990
A;Title: Nucleotide sequence analysis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       N;Alternate names: attachment glycoprotein G
C;Species: bovine respiratory syncytial virus
C;Date: 31-Mar-1992 #sequence_revision 31-Mar-1992 #text_change 16-Jul-1999
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A; Cross-references:
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-references: SGD:S0002553; MIPS:YDR146c
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Similarity 53.8%;
7; Conservation
61/1; 100/1; 151/2; 182/2;
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                                                                                             EMBL:AL049711; GSPDB:GN00061; ATSP:F4F15.320
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77.8%;
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Pred. No. 22;
1; Mismatches
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Pred. No.
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base, April 2000
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  222/2; 252/1; 297/3
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22;
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Matches 7; Conservative 2; Mismatches 5; Indels 0; Gaps 0;

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Search completed: January 29, 2003, 13:23:59

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1 15-JUN-2002 (Rel. 41, Last sequence update; 15-JUN-2002 (Rel. 41, Last annotation update)

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Trends Endocrinol. Metab. 12:11:
-!- FUNCTION: Specific ligand for
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                                                                             adiposity and stimulates gastric acid secretion. growth regulation.
SUBCELLULAR LOCATION: Secreted.
ALTERNATIVE PRODUCTS: 2 isoforms; 1/ghrelin (show 2/des-Gln14-ghrelin; are produced by alternative PTW: O-n-octanoylation is essential for activity.
SIMILARITY: BELONGS TO THE MOTILIN FAMILY.
DATABASE: NAME-Atlas Genet. Cytogenet. Oncol. Haer WWM: "http://www.infchiogenet.org/
                European
                                                                     WWW-"http://www.infobiogen.fr/services/chromcancer/Genes/GhrelinID.html"
                                                                                                                                                                                 FUNCTION: Specific ligand for the growth hormone secretagogue receptor type 1 (GHSR) inducing the release of growth hormone the pitultary. Has an appetite-stimulating effect, induces adiposity and stimulates gastric acid secretion. Involved in
                          SWISS-PROT entry is copyright. It is produced through a collaboration een the Swiss Institute of Bioinformatics and the EMBL outstation -
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2006/959; PubMed=10604470;
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SEQUENCE
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                                                                                                                             STRAIN=C57BL/6J;
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Staub A., Alexander G., Chenard M.-P., F
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Gastroenterology 119:395-405 (2000).
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                                                                                                                                       FROM N.A. (ISOFORM 1)
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, Hayashida
(APR-2001)
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51
117
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Rodentia;
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REMOVED IN MATURE FOR N-OCTANOATE.
MISSING (IN ISOFORM 2
                                                                                                                                                             ichi T., Nakao N.,
EMBL/GenBank/DDBJ
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Pred. No.
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Sciurognathi;
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secretagogue) (Growth
ed peptide) (M46 protei
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Rio M.-C.;
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thi; Muridae;
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databases.
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RESULT 4
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Best Local
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Q9QYH7; Q9ET69;
15-JUN-2002 (Rel. 4
15-JUN-2002 (Rel. 4
15-JUN-2002 (Rel. 4
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[5]
REVIEW.
MEDLINE=21203998;
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LIPID
VARSPLIC
SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This SWISS-PROT entry is copyright. It is produced through a collab between the Swiss Institute of Bioinformatics and the EMEL outst the European Bioinformatics Institute. There are no restrictions use by non-profit institutions as long as its content is in modified and this statement is not removed. Usage by and for comentities requires a license agreement (See http://www.isb-sib.ch/an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Brownstein M.J., Bult C., Fletcher C., Fujita M., Gariboldi M., Gustincich S., Hill D., Hofmann M., Hume D.A., Kamiya M., Lee N.H., Lyons P., Marchionni L., Mashima J., Mazzarelli J., Mombaerts P., Nordone P., Ring B., Ringwald M., Rodriguez I., Sakamoto N., Sasaki H., Sato K., Schoenbach C., Seya T., Shibata Y., Storch K.-F., Suzuki H., Toyo-oka K., Wang K.H., Weitz C., Whittaker C., Wilming L., Wynshaw-Boris A., Yoshida K., Hasegawa Y., Kawaji H., Kohtsuki S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Kojima M., Hosoda H., Matsuo H., Kangawa K.; "Ghrelin: discovery of the natural endogenous
                Ghrelin precursor
                                                                                                                                                                                                                                                                                                                                                       SIGNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         hormone secretagogue receptor.";
Trends Endocrinol. Metab. 12:118-122(2001).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       growth regulation.

SUBCELLULAR LOCATION: Secreted.

ALTERNATIVE PRODUCTS: 2 isoforms; 1/ghrelin (shown here) and ALTERNATIVE PRODUCTS: 2 isoforms; 1/ghrelin; esplicing.

2/des-Gln14-ghrelin; are produced by alternative splicing.

TISSUE SPECIFICITY: Mainly expressed in the gastrointestinal tracwith higher levels in the stomach, medium levels in the duodenum, jejunum, ileum and colon. Low expression in the testis and brain.

The description of the salivary gland, pancreas, liver and lung.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the pituitary. Has an appetite-stimulating effect, induces adiposity and stimulates gastric acid secretion formulation growth regulation.
                                                                                                                                                                                           GSSFLSPE----
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one; Cleavage
                                                                                                                                                            GSSFLSPEHQKAQQRKESKKPPAKLQPR
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AB060078; I
AK008658; I
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409:685-690(2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 detected in the salivary gland, pancreas, liver and lung.
: O-n-octanoylation is essential for activity (By similarity).
ILARITY: BELONGS TO THE MOTILIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   g requires a license agreement (S)
an email to license@isb-sib.ch).
                                                                                                                                                                                                                                    Similarity
peptide)
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                                                                                        STANDARD;
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                         41, Created)
41, Last sequence update)
41, Last annotation updat
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               (Growth hormone
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13207
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50.0%;
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REMOVED IN MATURE FORM (BY :
N-OCTANOATE (BY SIMILARITY)
MISSING (IN ISOFORM 2).
                                                                                                                                                                                                                                                                                                                                                                                 basic
                                                                                                                                                                                                                                  Score 53; I
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             secretagogue)
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                            update)
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1.012;
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               (Growth
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               hormone
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        a collaboration
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SEQUENCE FROM N.A. (ISOFORMS 1 AND 2), SEQUE SPECTROMETRY, AND ACYLATION OF SER-26. STRAIN=Sprague-Dawley; TISSUE=Stomach; MEDLINE=20357315; PubMed=10801861; MEDLINE=20357315, PubMed=10801861; MEDLINE=20357316, M., Matsuo H., Kangawa K., "Purification and characterization of rat deendogenous ligand for the growth hormone sec J. Biol. Chem. 275:21995-22000(2000).
                                                                                                                                                                                                                                                                                                                                                    "Ghreiin: qiscovery company (Ghreiin: qiscovery company (Ghreiin: qiscovery company (Ghreiin: qiscover)).

Trends Endocrinol. Metab. 12:118-122(2001).

Trends Endocrinol. Metab. 12:118-122(2001).

FUNCTION: Specific ligand for the growth hormone secretagogue receptor type 1 (GHSR) inducing the release of growth hormone the pituitary. Has an appetite-stimulating effect, induces the pituitary. Has an appetite acid secretion. Involved in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AND ACYLATION OF SER-26. SEQUENCE FROM N.A. (ISOFUKU A., ---
AND ACYLATION OF SER-26. STORMACH.
                                                                                                     This SWI
between
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MEDLINE-20067959; PubMed=10604470;
Kojima M., Hosoda H., Date Y., Nak
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=21433488; PubMed=11549267;
Matsumoto M., Hosoda H., Kitajima Y., Morozumi N.,
Tanaka S., Matsuo H., Kojima M., Hayashi Y., Kangaw
"Structure-activity relationship of ghrelin: pharma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                       modified and this statement is not removed.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kojima M., Hosoda H., Matsuo H., Ka
"Ghrelin: discovery of the natural
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Biochem.
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HOSOda H., Kojima M., Matsuo H., Kangawa
"Ghrelin and des-acyl ghrelin: two major
in gastrointestinal tissue.";
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                                                                                                                                                                                                    growth regulation.
SUBCELLULAR LOCATION: Secreted.
ALTERNATIVE PRODUCTS: 2 isoforms; 1/ghrelin (shown here) and ALTERNATIVE PRODUCTS: 2 isoforms; 1/ghrelin (shown here) and 2/des-Gln14-ghrelin; are produced by alternative splicing.
TISSUE SPECIFICITY: Broadly expressed with higher expression in the stomach. Very low levels are detected in the hypothalamus, heart, lung, pancreas, intestine and adipose tissue.
PTM: O-n-octanoylation is essential for activity. The replacement of Ser-26 by aromatic tryptophan preserves ghrelin activity.
MASS SPECTROMETRY: MW=3314.9; MW_ERR=0.7; METHOD=Electrospray;
                                                                     s SWISS-PROT entry is copyright. It is produced through a cween the Swiss Institute of Bioinformatics and the EMBL European Bioinformatics Institute. There are no restrict European Bioinformatics Institute. There are no restrict by non-profit institutions as long as its content is
                                                                                                                                                                            MASS SPECTROMETRY:
                                                                                                                                                                 RANGE=24-36,
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                          an
                                         requires a license agreement
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                          email to license@isb-sib.ch).
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Da; Chordata;
la; Rodentia;
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                                                                                                                                                                               MW=3187.1;
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Sciurognathi;
                                                                                                                                                 MOTILIN FAMILY.
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t des-Gln14-ghrelin,
secretagogue recept
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                                                       Usage
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pharmacological study
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15-JUN-2002 (Rel. 41, Created)
15-JUN-2002 (Rel. 41, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation updat
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                                                                                                                                                                                                                                                                                                                   the European Bioinformatics Institute. The by non-profit institutions as long
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Submitted (MAR-2001)
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Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                                                                                                                                                            growth regulation (By similarity).
SUBCELLULAR LOCATION: Secreted (By similarity).
ALTERNATIVE PRODUCTS: 2 isoforms; 1/ghrelin (shown here) and 2/des-Gln14-ghrelin; are produced by alternative splicing.
PTM: O-n-octanoylation is essential for activity (By similarity).
SIMILARITY: BELONGS TO THE MOTILIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         mitted (MAR-2001) to the EMBL/GenBánk/DDBJ databases. FUNCTION: Specific ligand for the growth hormone secretagogue receptor type 1 (GHSR) inducing the release of growth hormone the pituitary. Has an appetite-stimulating effect, induces adiposity and stimulates gastric acid secretion. Involved in
                                                                                                                                                                             AB035704; BAB19049.1; -. AF308930; AAK19243.1; -.
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SEQUENCE
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Submitted (JAN-1994) to the EMBL/GenBank/DDBJ databases.
-i- FUNCTION; UNLIKE THE OTHER PARAMYXOVIRUS ATTACHMENT PR
RESPIRATORY SYNCYTIAL VIRUS G PROTEIN LACKS BOTH NEURA
HEWAGGLUTINATING ACTIVITIES.
-i- SUBCELLULAR LOCATION: EXPRESSED ON THE SURFACE OF THE
CELLS AND INCORPORATED IN THE MEMBRANE OF THE VIRIONS.
-i- SIMILARITY: TO THE G PROTEINS OF EITHER THE SUBGROUP A
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Pfam; PF00802; Glycoprotein G; 1.
Transmembrane; Glycoprotein.
DOMAIN 1 37
                                                                                                                                                                                                                  EMBL; L10925; AAA42809.1;
HSSP; P22261; 1BRV.
                                                                                                                                                                                                                                               or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Viruses; ssRNA negative-strand viruses; Mononegavirales; Paramyxoviridae; Pneumovirinae; Pneumovirus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      respiratory syncytial virus G glycoprotein.";
J. Gen. virol. 78:359-366(1997)
-I- FUNCTION: UNLIKE THE OTHER PARAMYXOVIRUS ATTACHMENT PROTEINS, 7:
RESPIRATORY SYNCYTIAL VIRUS G PROTEIN LACKS BOTH NEURAMINIDASE
HEMAGGLUTINATING ACTIVITIES.
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MEDLINE=97170754; PubMed=9018058; .
Walravens K., Langedijk J.P.,
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         Bovine respiratory syncytial virus (strain Rb94) (BRS), Bovine respiratory syncytial virus (strain 220-69) (BRS) Viruses; serna negative-strand viruses; Mononegavirales;
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(Rel. 36, Last annotation update)
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J. Gen. Virol. 78:359-366(1997).
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Letesson J.J.;
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Kojima M.
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STRAIN=ATCC 29342
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Bacteria; Firmicutes; Mollicutes; Mycoplasmataceae; Mycoplasma
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                                    -1- SIMILARITY: BELONGS TO THE S18P FAMILY OF RIBOSOMAL PROTEINS
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-1- FUNCTION: Specif:
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SUBCELLULAR LOCATION: Secreted (By similarity).
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Best Local S
Matches 8
        InterPro; IFR001648; Ribosomal S18.

Pfam; PF01084; Ribosomal S18; T.

PRINTS; PR00974; RIBOSOMALS18.

ProDom; PD002239; Ribosomal S18; 1.

TIGR002239; Ribosomal S18; FALSE NEG.

PROSITE; PS00057; RIBOSOMAL S18; FALSE NEG.

RIBOSOMAL S18; Complete per Ribosomal protein; RNA-binding; Complete per Ribosomal per Ribos
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P47338;
01-FEB-1996
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InterPro; IPR001648; Ribosomal S18; T.
Pfam; PF01084; Ribosomal S18; T.
PRINTS; PR00974; RIBOSOWALS18.
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TIGRPAMs; TIGR00165; S18; T.
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SEQUENCE 104 AA; 12386 MW; 0C424229C79CC1B5 CR
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P08153;
01-AUG-1988
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Q9J552;
16-OCT-2001
16-OCT-2001
16-OCT-2001
SEQUENCE FROM N.A.
MEDLINB=88211561; PubMed=3284746;
Stillman D.J., Bankier A.T., Seddon A
"Characterization of a transcription
specific transcription of the yeast H
EMBO J. 7:485-494(1988).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This SWISS-PROT entry is copyright. It is produced through a collable between the Swiss Institute of Bioinformatics and the EMBL outsithe European Bioinformatics Institute. There are no restrictions use by non-profit institutions as long as its content is in modified and this statement is not removed. Usage by and for conmodified and this statement is not removed.
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MEDLINE=20193820; PubMed=10729156;
MEDLINE=20193820; FubMed=10729156;
                                                                                                                                                                            Saccharomyces cerevisiae (Baker's yeast).
Eukaryota; Fungi; Ascomycotta; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetacese; Saccharomyces.
                                                                                                                                                                                                                                           Transcriptional factor SWI5. SWI5 OR YDR146C OR YD8358.03C.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Afonso C.L., Tulman E.R., Lu Z., Zsak L.,
"The genome of fowlpox virus.";
J. Virol. 74.3815-8831(2000).
-i- SIMILARITY: BELONGS TO THE POXVIRUSES
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Viruses; dsDNA viruses,
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UR InterPro; IPR000637; AT hook.

DR InterPro; IPR000822; ZnF C2H2.

DR Pfam; PP00096; zf-C2H2; 3.

DR Pfam; PP00178; AT hook; 1.

DR PRINTS; PR00048; ZINCFINGER.

DR SMART; SM00384; AT hook; 1.

DR SMART; SM00385; ZnF C2H2; 2.

DR PROSITE; PS00028; ZINC FINGER C2H2 1; 2.

DR PROSITE; PS00028; ZINC FINGER C2H2 2; 2.

VW Transcription regulation; Activator; Zinc-finger; DNA-binding;

VW Repeat; Metal-binding; Nuclear protein; Phosphorylation;

VW 3D-structure.
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EMBL; Z50046; CAA90369.1; -
PIR; S00342; TWBYS5.
PDB; 1NCS; 10-JUL-96.
PDB; 1ZFD; 14-OCT-96.
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SEQUENCE FROM N.A.
STRAIN=S288C / AB972,
Murphy L., Richards (
Walsh S.V.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This SWISS-PROT entry is copyright. It is produced through a collab between the Swiss Institute of Bioinformatics and the EMBL outst the European Bioinformatics Institute. There are no restrictions use by non-profit institutions as long as its content is in use by non-profit institutions as long as its content of content is in the content of the content in the content is not removed. Usage by and for content is not removed.
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Neolution structures of two zinc-finger domains from SWI5 obtained using two-dimensional 1H nuclear magnetic resonance spectroscopy. A zinc-finger structure with a third strand of beta-sheet.";

J. Mol. Biol. 228:637-651(1992).

-!- FUNCTION: DETERMINES THE MOTHER-CELL-SPECIFIC TRANSCRIPTION OF HO ENDONUCLEASE GENE THAT IS RESPONSIBLE FOR THE INITIATION OF MATING-TYPE SWITCHING IN YEAST. RECOGNIZES A SPECIFIC SEQUENCE THE PROMOTER OF THE HO GENE. ACTIVATES EGT2 TRANSCRIPTION IN A CONCENTRATION-DEPENDENT MANNER. SYNTHESIZED DURING G2 AND EARLY CONCENTRATION-DEPENDENT MANNER.
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factor SWI5.";
Cell 66:743-758(1991).
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Neuhaus D., Nakase
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"Zinc-finger motifs expressed
specific binding to DNA.";
Nature 332:284-286(1988).
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Nagai K., Nakaseko
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SUBCELLULAR LOCATION: NUCLEAR IN G1, BUT CYTOPLASMIC IN 5, G4, MITOSIS.

MICCLI CYCLE PHASES:
MICCLI CYCLE-DEPENDENT PHOSPHORYLATION OF THREE SERINE RESIDUES PREVENTS SWI5 FROM ENTERING THE NUCLEUS, AND IT ACCUMULATES IN THE CYTOPLASM. AS A CONSEQUENCE OF CDC28 KINASE INACTIVATION AT THE END OF ANAPHASE, THE THREE SERINE RESIDUES ARE DEPHOSPHORYLATED AND SWI5 ENTERS THE NUCLEUS TO ACTIVATE TRANSCRIPTION. IT IS THEN RAPIDLY DEGRADED. THREONINE PHOSPHORYLATION ALSO SEEMS TO OCCUR. RAPIDLY STRONG, TO YEAST METALLOTHIONEIN EXPRESSION ACTIVATOR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 an email to license@isb-sib.ch).
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O Y., Nasmyth K.A.,
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cerevisiae transcription
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Best Local S
Matches 7
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JUNG K.-H., Christensen D.J., Scott A.I.;
"Isolation and characterization of two genes from Camptotheca acuminata that encode isopentenyl diphosphate isomerase.";
Submitted (OCT-1997) to the EMBL/GenBank/DBU databases.
-i-FUNCTION: CATALYZES THE 1.3-ALLYLIC REARRANGEMENT OF THE HOMOALLYLIC SUBSTRATE ISOPENTENYL (IPP) TO ITS ALLYLIC ISOME DIMETRYLALLYL DIPHOSPHATE (DMAPP) (BY SIMILARITY).
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Asteridae; Cornales; Cornaceae; Nyssoideae; Camptotheca.
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MOD_RES
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InterPro; IPR000086; NUDIX_hydrolase.
Pfam; PF00293; NUDIX; 1.
                                                                                                                                                                                                                            entities requires a license agreement (S or send an email to license@isb-sib.ch).
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30-MAY-2000 (Rel. 39, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Isopentenyl-diphosphate delta-isomerase II (E
II) (Isopentenyl pyrophosphate isomerase II).
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30-MAY-2000
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RESULT 15
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01-AUG-1991 (Rel. 1
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PDB; 1BRV;
                                                                                    DOMAIN
DISULFID
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Biochemistry 35:14684-14688 (1996).

-i- FUNCTION: UNLIKE THE OTHER PARAMYXOVIRUS ATTACHMENT EN RESPIRATORY SYNCYTIAL VIRUS G PROTEIN LACKS BOTH NEURI-HEMAGGLUTINATING ACTIVITIES.

-i- SUBCELLULAR LOCATION: EXPRESSED ON THE SURFACE OF THE CELLS AND INCORPORATED IN THE MEMBRANE OF THE VIRIONS.

-i- SUBCELLULARITY: TO THE G PROTEINS OF EITHER THE SUBGROUP I
                                                                                                                                                        Pfam;
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 SEQUENCE
                                                                                                                                                                                                                                          entities requires a license agreement (S or send an email to license@isb-sib.ch).
                                                                                                                TRANSMEM
                                                                                                                             DOMAIN
                                                                                                                                           Transmembrane;
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Rullmann J.A., Schaaper W.M., van Oirschot
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    syncytial virus is distinct from
virus.";
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A36408; MGNZBR
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Pred. No. 9.2;
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                                                                                                                             (POTENTIAL).
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| Search completed: January 29, 2003, 13:23:01 Job time : 13 secs | Query Match Best Local Similarity 77.8%; Pred. No. 9.4; Matches 7; Conservative 1; Mismatches 1; Indels 0; Gaps 0; Qy 6 SPEAKLQPR 14 |
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Listing first 45 s
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         111111 98 7654321
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1: sp_archea:*

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Q8tat9 homo sapien P89309 bovine resp P90378 bovine resp P89310 bovine resp Q9ynf9 bovine resp Q9ynf8 bovine resp Q9ynf6 bovine resp Q9ynf5 caenorhabdi Q9dbf6 mus musculu Q47732 enterococcu Q98f99 rhizobium 1 Q9aab3 caulobacter
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|--------------------|--------------------|--------|--------------------|-------------------|--------|--------|--------------------|--------|--------------------|-------------------|--------------|--------------------|--------------------|----------|--------------------|--------------------|--------|--------------------|--------------------|--------------------|-------------------|--------------------|-------------------|--------------------|--------|--------|--------------------|--------------------|
| 54.3 | 54.3 | | 54.3 | | | 54.3 | 55.0 | 55.7 | 55.7 | 55.7 | 55.7 | 55.7 | 55.7 | 55.7 | 55.7 | 55.7 | 55.7 | 55.7 | 55.7 | 55.7 | 57.1 | 57.1 | 57.1 | 57.1 | 57.1 | 57.1 | 57.1 | 57.1 |
| 459 | 406 | 322 | 226 | 187 | 108 | 103 | 681 | 1336 | 1335 | 991 | 558 | 558 | 532 | 532 | 473 | 469 | 435 | 371 | 251 | 188 | 1078 | 680 | 470 | 343 | 343 | 308 | 256 | 252 |
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| 800660 | Q9P8D1 | 045024 | Q9RT86 | Q970U3 | Q40737 | Q8VDZ2 | 081436 | Q9ES55 | Q8VI17 | Q9VIF5 | Q9R1W1 | Q9WTL5 | Q9WTL6 | Q9R1W2 | Q9ZC65 | Q9CS26 | Q8R162 | Q9SUY4 | P89311 | Q92KN0 | Q9NDY3 | Q921G6 | Q8X0Z2 | Q8YAM9 | Q92FH1 | Q9AAX7 | Q9S0A7 | P70866 |
| Q99u08 staphylococ | Q9p8d1 cephalospor | | Q9rt86 deinococcus | Q970u3 sulfolobus | | | O81436 arabidopsis | | Q8vi17 mus musculu | Q9vif5 drosophila | \mathbf{r} | Q9wtl5 mus musculu | Q9wtl6 mus musculu | mus musc | Q9zc65 yersinia pe | Q9cs26 mus musculu | | Q9suy4 arabidopsis | P89311 bovine resp | Q92kn0 rhizobium m | Q9ndy3 leishmania | Q921g6 mus musculu | Q8x0z2 gibberella | Q8yam9 listeria mo | | 7 | Q9s0a7 borrelia bu | P70866 borrelia bu |

ALIGNMENTS

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RESULT
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Best Local (
                                       P89309;
P89309;
01-MAY-1997
01-MAY-1997
01-MAR-2002
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Q8TAT9;
01-JUN-2002
01-JUN-2002
                                                                                                                                                                                                                                                                     Strausberg R.;
Submitted (MAR-2002) to the EMBL/GenBank/DDBJ databases.
EMBL; BC055791; AAH25791.1; -.
SEQUENCE 117 AA; 12929 MW; 25B0572EBECB7610 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens (Human)
Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=9606; [1]
                                                                                                                                                                                                                                                                                                                                                                                                                       Ghrelin.
Bovine respiratory syncytial virus
                          Attachment glycoprotein
                                                                                                                                                                                                                                                                                                                            TISSUE-BLOOD;
                                                                                                                                                                                                                                                                                                                                                                                                                                       01-JUN-2002
                                                                                                                                                                                                                                                                                                                                         EQUENCE FROM N.A.
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                                                                                                                                                             14;
                                                                                                                                                                                                                              h 75.7%;
Similarity 50.0%;
               (TYEMBLrel. 03, Created)
(TYEMBLrel. 03, Last sequence update)
(TYEMBLrel. 20, Last annotation update)
clycoprotein (Fragment).
                                                                                                                                                                                                                   Conservative
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(TrEMBLrel. 21, Last sequence update)
(TrEMBLrel. 21, Last annotation update)
                                                                                              PRELIMINARY;
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Primates;
                                                                                                                                                                                                                 0; Mismatches
                                                                                                                                                                                                                               Score 53; DB 4; Length 117; Pred. No. 0.031;
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                                                                                              251 AA
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P90378

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Best Local S
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J. Gen. Virol. 78:359-366(1997)

EMBL; U24714; AACS6941.1; -.

HSSP; P22261; 1BRV.

InterPro; IPR001925; Glycoprot G.

InterPro; IPR001880; Ppantne attach.
Pfam; PF00802; Glycoprotein G; 1.

PROSITE; PS00012; PHOSPHOPANTETHEINE; UNKNOWN 1.
                                                                                                                                                                                                                                                                                                                                  PROSITE;
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P90378,
01-MAY-1997 (TrEMBLrel. 03, Created)
01-MAY-1997 (TrEMBLrel. 03, Last seguence update)
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
                                                                                                                                                                                                                                                                                                       SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Prozzi D., Walravens
Letesson J.J.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=97170754; PubMed=9018058;
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                                                                                                                       218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bovine respiratory syncytial virus.
Viruses; sgRNA negative-strand viruses; Mononegavirales;
Paramyxoviridae; Pneumovirinae; Pneumovirus.
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InterPro; IPR003880; Ppantne attach.
Pfam; PF00802; Glycoprotein G; 1.
PROSITE; PF008012; PHOSPHOPANTETHEINE; UNKNOWN_1.
NON_TER 1
1 1 CED605DD49103DC3
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EMBL; U24713; AAC56940.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Antigenic and molecular analyses of the variability of bovine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Attachment glycoprotein
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Letesson J.J.;
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Paramyxoviridae; Pneumovirinae; Pneumovirus.
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                                                                                                                                                                                                             8; Conserv
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                                                                                                                                                                                                                                                                                                          251 AA;
  PRELIMINARY;
                                                                                                                                                                                                          Conservative
                                                                                                                     226
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ens K., Langedijk
                                                                                                                                                                                                                                                                                                     27680 MW;
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88.9%;
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88.9%;
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Pred. No. 5.6;
1; Mismatches
                                                                                                                                                                                                   Score 43; DB 12;
Pred. No. 5.6;
1; Mismatches 0
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01-MAY-1997 (TrEMBLrel. 03
01-MAY-1997 (TrEMBLrel. 03
01-MAR-2002 (TrEMBLrel. 27
Attachment glycoprotein (
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Q9YNF9;
01-MAY-1999
01-MAY-1999
01-MAR-2002
                                                                                                                                   InterPro; IPR000925; Glycoprot G.
InterPro; IPR003880; Ppantne altach.
Pfam; PF00802; Glycoprotein G; I
PROSITE; PS00012; PHOSPHOPANTETHEINE; UNKNOWN 1.
SEQUENCE 257 AA; 28288 MW; C3106210BACA2817
                                                                                                                                                                                                                                                                                                                   Larsen L.E., Utenthal A., Arctander P., Tjornehoj K., Viuff B., Rontved C., Ronsholt L., Alexandersen S., Blixenkrone-Moller M.; "Serological and genetic characterization of bovine respiratory syncytial virus (BRSV) indicates that Danish isolates belong to the intermediate subgroup: no evidence of a selective effect on the variability of grotein nucleotide sequence by prior cell culture adaption and passages in cell culture or calves."; vet. Microbiol. 62:255-279(1998).

EMBL, U92098; ADD00710.1; -.

EMBL, U92098; ADD00710.1; -.
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Viruses; ssRNA negative-strand viruses; Mononegavirales;
Paramyxoviridae; Pneumovirinae; Pneumovirus.
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NCBI_TaxID=11246;
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Pfam; PF00802; Glycoprotein_G; 1.
PROSITE; PS00012; PHOSPHOPANTETHEINE; UNKNOWN_1.
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SEQUENCE 251 AA; 27643 MW; 7786571D0EFF9D51
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J. Gen. Virol. 78:359-366(1997).
EMBL; U24715; AAC556942.1; -.
HSSP; P22261; 1BRV.
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Viruses; ssRNA negative-strand viruses; Mononegavirales;
Paramyxoviridae; Pneumovirinae; Pneumovirus.
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Letesson J.J.;
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ens K., Langedijk J.P.,
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Larsen L.E., Uttenthal A., Arctander P., Tjornehoj K., Viuff B., Rontved C., Ronsholt L., Alexandersen S., Blixenkrone-Moller M., "Serological and genetic characterization of bovine respiratory syncytial virus (BRSV) indicates that Danish isolates belong to the intermediate subgroup: no evidence of a selective effect on the variability of G protein nucleotide sequence by prior cell culture adaption and passages in cell culture or calves."; vet. Microbiol. 62:265-279(1998).
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EMBL; U9210; AAD00712.1; -.

HSSP; P22261, 1BRV.
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InterPro; IPR003880; Ppantne attach.
Pfam; PP00802; Glycoprotesin G; 1:
PROSITE; PS00012; PHOSPHOPANTETHEINE; UNKNOWN 1.
SEQUENCE 257 AA; 28324 MW; 94EE8BBC358739B8 CRC64;
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Viruses; ssRNA negative-strand viruses; Mononegavirales;
Paramyxoviridae; Pneumovirinae; Pneumovirus.
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Viruses; ssRNA negative-strand viruses; Mononegavirales;
Paramyxoviridae; Pneumovirinae; Pneumovirus.
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24 SPEAKLQPK 232
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Q9YNF6;
01-MAY-1999
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EMBL; U92102; AAD00714.1; -.

HSSP; P22261; 1BRV.
Paramyxoviridae;
NCBI_TaxID=11246;
[1]
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InterPro; IPR003880; Ppantne attach.
Pfam; PF00802; Glycoprotein G; 1.
PROSITE; PS00012; PHOSPHODANTETHEINE; UNKNOWN 1.
SEQUENCE 257 AA; 28380 MW; 6BD146B3B02E5ACB
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                                                                Bovine respiratory syncytial virus.
Viruses; ssRNA negative-strand viruses; Mononegavirales;
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InterPro; IPR003880; Ppantne attach.
Pfam; PF00802; Glycoprotein G; 1.
PROSITE; PS00012; PHOSPHOPANTETHEINE; UNKNOWN 1.
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Viruses; ssRNA negative-strand viruses; Monc
Paramyxoviridae; Pneumovirinae; Pneumovirus
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01-MAR-2002
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                                           Pneumovirinae; Pneumovirus
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88.9%;
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RESULT 11
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WHEL W22104; AAD00716.1; -.
    Q9YS25
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InterPro; IPR003880; Ppantne attach.
Pfam; PF00802; Glycoprotein G; 1.
PROSITE; PS00012; PHOSPHOPANTETHEINE; UNKNOWN 1.
SEQUENCE 257 AA; 28379 MW; ClAZD366EC1657TE CRC64;
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Viruses; seRNA negative-strand viruses; Mononegavirales;
Paramyxoviridae; Pneumovirinae; Pneumovirus.
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01-MAR-2002 (TrEMBLrel.
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InterPro; IPRO00925; Glycoprot G.
InterPro; IPRO00926; Ppantne attach.
InterPro; IPRO03880; Ppantne attach.
Pfam; PP00802; Glycoprotein G; 1.
Pfam; PP00802; Glycoprotein G; 1.
PROSITE; PS00012; PHOSPHOPANTETHEINE; UNKNOWN 1.
PROSITE; PS00012; PHOSPHOPANTETHEINE; UNKNOWN 1.
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EMBL, U92103, AAD00715.1; -.

HSSP; P22261; 1BRV.
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STRAIN=9402020
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1; Mismatches
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RESULT 12
Q9XUV5
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OT 01-NO
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OC ENKAZ
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Q9XUV5;
01-NOV-1999
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01-DEC-2001
  SEQUENCE FR
Lloyd C.R.;
                                                           Rhabditidae; Pel
NCBI_TaxID=6239;
                                                                                                        Caenorhabditis elegans.
Eukaryota; Metazoa; Nem
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InterPro; IPR003880; Ppantne attach.
Pfam; PF00802; Glycoprotein G; 1.
PROSITE; PS00012; PHOSPHOPANTETHEINE;
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SPECIES-Bovine respiratory syncytial virus; STRAIN-ATUE51908;
MEDLINE-99102581; PubMed-9847328;
Buchholz U.J., Finke S., Conzelmann K.K.;
Buchholz U.J., Finke S., Strainson Strainson (BRSV) from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SPECIES=Bovine respiratory syncytial virus; STRAIN=ATUE51908; Buchholz U.J., Finke S., Conzelmann K.-K.; Submitted (SEP-1998) to the EMBL/GenBank/DDBJ databases.
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SPECIES=Bovine res
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01-MAY-1999 (TrEMBLrel.
01-JUN-2002 (TrEMBLrel.
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8; Conserv
                      FROM N.A
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                                                                                                                                                                                        (TrEMBLrel. (TrEMBLrel. (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     257
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                                                                               Peloderinae;
                                                                                                                                                                                                                                                                                  PRELIMINARY;
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                                                                                                   Nematoda; Chromadorea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 61.4%;
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                                                                                    Caenorhabditis.
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Last ann
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Pred. No.
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annotation update)
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                                                                                                      Rhabditida;
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Query Match
Best Local Similarity
Watches B; Conserve
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RA Arakawa T., Hara A., Fukunishi Y., Konno H., Adachi J., Fukuda S.,
RA Aizawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I.,
RA Aizawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I.,
RA Aizawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I.,
RA Saito T., Okazaki Y., Gojobori T., Bono H., Kasukawa T., Saito R.,
RA Kadota K., Matsuda H.A., Ashburner M., Bacalov S., Casavant T.,
RA Kadota K., Matsuda H.A., Ashburner M., Bacalov S., Casavant T.,
RA Kadota K., Matsud Y., Nikaido I., Pesole G., Quackenbush J.,
RA Kuehl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Kuehl P., Lewis S., Matsuo Y., Nikaido I., Pesole G., Quackenbush J.,
RA Schriml L.M., Staubli F., Suzuki R., Tomita M., Wagner L., Washio T.,
RA Schriml L.M., Staubli F., Suzuki R., Tomita M., Wagner L., Washio T.,
RA Sakai K., Okido T., Furuno M., Aono H., Baldarelli R., Barsh G.,
RA Blake J., Boffelli D., Bojunga N., Carninci P., de Bonaldo M.F.,
RA Blake J., Boffelli D., Hofmann M., Hume D.A., Kamiya M., Lee N.H.,
RA Gustincich S., Hill D., Hofmann M., Hume D.A., Kamiya M., Lee N.H.,
RA Lyons P., Marchionni L., Mashina J., Mazzarelli J., Mombaerts P.,
RA Sasaki H., Sato K., Schoenbach C., Seya T., Shibata Y., Storch K.-F.,
RA Mynshaw-Boris A., Yoshida K., Hasegawa Y., Kawaji H., Kohtsuki S.,
RA Hayashizaki Y.;
  Query Match
Best Local S
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
STRAIN=C57BL/6J; TISSUE=LUNG;
MEDLINE=21085660; PubMed=11217851;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            investigating biology,"; science 282:2012-2018(1998). EMBL; 281560; CAB04544.1; -. SEQUENCE 164 AA; 19025 MW;
                                                                                                                                PRINTS; PRO0401; SH2DOMAIN
ProDom; PD000093; SH2; 1.
SMART; SM00462; PTB; 1.
SMART; SM00252; SH2; 1.
PROSITE; PS50001; SH2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Q9DBT6;
01-JUN-2001 (TrEMBLrel.
01-JUN-2001 (TrEMBLrel.
01-MAR-2002 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1200014E20Rik protein.
TNS OR 1200014E20RIK.
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                                                                                                             SEQUENCE
                                                                                                                                                                                                                                 InterPro; IPR000050; PID domain.
InterPro; IPR000980; SH2.
Pfam; PF00017; SH2; 1.
PRINTS; PR00401; SH2DOMAIN.
                                                                                                                                                                                                                                                                                                                                                                                               EMBL; AK004758; BAB23539.1;
                                                                                                                                                                                                                                                                                                                                                                                                                    "Functional annotation of a full-length mouse cDNA collection."; Nature 409:685-690(2001).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mus musculus (Mouse).
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MEDLINE=99069613; PubMed=9851916;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=10090;
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                                                                                                                                                                                                                                                                                                                                            MGD; MGI:104552; Tns.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              13
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                                                                                                                                                                                                                                                                                                                                                                 P16277; 1BLK
     Similarity
8; Conser
  58.6%;
nilarity 61.5%;
Conservative ;
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                                                                                                        624 AA;
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                                                                                                           65369 MW;
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  Score 41; DB:
Pred. No. 36;
2; Mismatches
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Last annotation update)
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Pred. No. 8.4;
2; Mismatches
     <u>ب</u>
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                                                                                                           296FC3741EFBC57E CRC64;
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                                                     Length 624;
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Best Local S
Matches 8
                                                                                Matches
                                                                                                                                Query Match
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Q98F99;
01-OCT-2001
01-OCT-2001
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Q47732;
01-NOV-1996
01-NOV-1996
                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.

STRAIN-MAFF303099;

MEDLINE-21082930; PubMed=11214968;

Kaneko T., Nakamura Y., Sato S., Asamizu E., Kato T., Sasa Watanabe A., Idesawa K., Ishikawa A., Kawashima K., Kimura Kishida Y., Kiyokawa C., Kohara M., Matsumoto M., Matsuno Mochizuki Y., Nakayama S., Nakazaki N., Shimpo S., Sugimot Takeuchi C., Yamada M., Tabata S.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Flannagan S.E., Zitzow L.A., Su Y.A., "Nucleotide sequence of the 18-kb con Enterococcus faecalis.";
Plasmid 32:350-354(1994).
EMBL; U09422; AAB60017.1; -.
SEQUENCE 815 AA; 93317 MW; 594C6A:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hypothetical MSR3871.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRAIN=DS16; TRANSPOSON=TN916;
MEDLINE=95207419; PubMed=7899523;
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Enterococcaceae; Enter
                                                                                                                                                                                                        EMBL; AP003002; BAB50668.1; Hypothetical protein; Comple
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-MAR-2002
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                                                                                                                                                                                   SEQUENCE
                                                                                                                                                                                                                                                           Mesorhizobium loti.";
DNA Res. 7:331-338(2000).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Phyllobacteriaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bacteria; Proteobacteria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Rhizobium loti (Mesorhizobium loti)
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                                                                                                                                                                                                                                                                                                              'Complete genome structure
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15
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1 GSSFLSPEAKLQ
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8; Conserv
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8; Conserv
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(TrIMBLrel. 18, Last sequence up
(TrIMBLel. 20, Last annotation
protein msr3871.
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                                                                                Conservative
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6 AA; 10602
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